## SEQUENCE LISTING

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Lys Pro Thr 

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Glu Ile Asp Ser Val Gly Ser Glu Asp Ser Leu Asp Thr Ser Leu Arg 50 55 60

Ala His Gly Val His Ala Thr Lys His Val Pro Glu Lys Arg Pro Leu 65 70 75 80

Pro Ile Arg Arg Lys Arg Ser Ile Glu Glu Ala Val Pro Ala Val Cys 85 90 95

Lys Thr Arg Thr Val Ile Tyr Glu Ile Pro Arg Ser Gln Val Asp Pro
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Thr Ser Ala Asn Phe Leu Ile Trp Pro Pro Cys Val Glu Val Lys Arg 120 125 115 Cys Thr Gly Cys Cys Asn Thr Ser Ser Val Lys Cys Gln Pro Ser Arg 135 140 130 Val His His Arg Ser Val Lys Val Ala Lys Val Glu Tyr Val Arg Lys Lys Pro Lys Leu Lys Glu Val Gln Val Arg Leu Glu Glu His Leu Glu 165 170 Cys Ala Cys Ala Thr Thr Ser Leu Asn Pro Asp Tyr Arg Glu Glu Asp 190 185 Thr Asp Val Arg 195 <210> 5 <211> 6633 <212> DNA <213> Homo sapiens <220> <221> CDS <222> (395)..(3661) <400> 5 ttctccccgc ccccagttg ttgtcgaagt ctgggggttg ggactggacc ccctgattgc 60 gtaagagcaa aaagcgaagg cgcaatctgg acactgggag attcggagcg cagggagttt 120 gagagaaact tttattttga agagaccaag gttgaggggg ggcttatttc ctgacagcta 180 tttacttaga gcaaatgatt agttttagaa ggatggacta taacattgaa tcaattacaa 240 aacgcggttt ttgagcccat tactgttgga gctacaggga gagaaacagg aggagactgc 300 aagagatcat ttgggaaggc cgtgggcacg ctctttactc catgtgtggg acattcattg 360 cggaataaca tcggaggaga agtttcccag agct atg ggg act tcc cat ccg gcg 415 Met Gly Thr Ser His Pro Ala 463 ttc ctg gtc tta ggc tgt ctt ctc aca ggg ctg agc cta atc ctc tgc Phe Leu Val Leu Gly Cys Leu Leu Thr Gly Leu Ser Leu Ile Leu Cys 15 10 cag ctt tca tta ccc tct atc ctt cca aat gaa aat gaa aag gtt gtg 511 Gln Leu Ser Leu Pro Ser Ile Leu Pro Asn Glu Asn Glu Lys Val Val

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Phe Val Thr Val Leu Glu Val Ser Ser Ala Ser Ala Ala His Thr Gly 85 90 95

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Phe Ser Gln Leu Glu Ala Val Asn Leu His Glu Val Lys His Phe Val 325 330 335

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Glu Gln Arg Leu Ser Ala Asp Ser Gly Tyr Ile Ile Pro Leu Pro

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Ser Ala Ile Ile Pro Cys Arg Thr Thr Asp Pro Glu Thr Pro Val Thr 150 155 145 Leu His Asn Ser Glu Gly Val Val Pro Ala Ser Tyr Asp Ser Arg Gln

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135

170

95

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Ile Cys Lys Asp Ile Lys Lys Cys Asn Asn Glu Thr Ser Trp Thr Ile 450 455 460

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